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1 : Immunology 1993 Jan;78(1):65-73

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## Characterization of T-cell responses to the house dust mite aller Der p II in mice. Evidence for major and cryptic epitopes.

Hoyne GF, Callow MG, Kuo MC, Thomas WR

Western Australian Research Institute for Child Health, Princess Margaret Hospital for Children, Perth.

Major histocompatibility complex (MHC) congenic strains can be defined a and low responders to the major house dust mite allergen Der p II on the ba the ability to sensitize T cells for in vitro lymphokine release. Mice of the H haplotype were high responders, H-2k were intermediate and H-2d low responders. Like responses to other proteins, only a limited number of epito could be located by the response of T cells from mice immunized with aller a series of overlapping peptides. The epitopes for H-2b mice were 11-35, 78 and 105-129, 36-50 and 78-104 for H-2k mice and 36-60 for H-2d. Immuni with the peptides however revealed that spleen-adherent cells were required lymph node cells to recall responses to the whole protein and in addition tha could be sensitized by cryptic epitopes defined by peptides 22-50 and 1-20 f 2b mice. Peptides containing these cryptic epitopes did not normally induce responses in mice primed with the allergen, but when they were used for immunizing they could prime mice for responses to the peptide and the who allergen. The results both help to define a model for studying the presentatio allergens and have significant implications for peptide-based immunotherap

PMID: 7679663, UI: 93170864

☒

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**Nucleotide**

BLAST PubMed Nucleotide Protein Genome Structure PopSet Taxonomy Help  
Sequence 79 from patent US 5776761

**Views:**

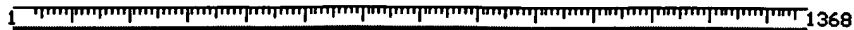
Accession: AR016640  
Total Bases Sequenced: 1368 bp  
Completed: Dec 5, 1998.

**GenBank view**

US PATENT: 5776761

FASTA view

### ASN.1 view



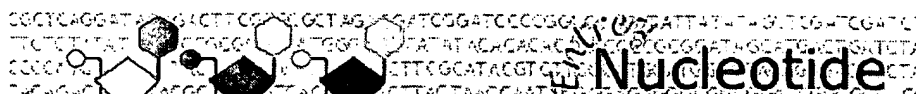
**Legend:**

██████ - 100-200-...

Unknown. Unclassified.

**Nucleic acids encoding allergenic proteins from ragweed**  
Rogers,B., Klapper,D.G., Rafnar,T. and Kuo,M.  
Patent: US 5776761-A 79 07-JUL-1998;

*Comments and suggestions to: [info@ncbi.nlm.nih.gov]*



Search  for

1 : [AR016640](#) . Sequence 79 from p...[gi:3972917]

Related Sequences

LOCUS AR016640 1368 bp DNA PAT 05-DEC-1998  
 DEFINITION Sequence 79 from patent US 5776761.  
 ACCESSION AR016640  
 VERSION AR016640.1 GI:3972917  
 KEYWORDS .  
 SOURCE Unknown.  
 ORGANISM Unknown.  
 UNCLASSIFIED.  
 REFERENCE 1 (bases 1 to 1368)  
 AUTHORS Rogers,B., Klapper,D.G., Rafnar,T. and Kuo,M.  
 TITLE Nucleic acids encoding allergenic proteins from ragweed  
 JOURNAL Patent: US 5776761-A 79 07-JUL-1998;  
 FEATURES Location/Qualifiers  
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 /organism="unknown"  
 BASE COUNT 397 a 286 c 327 g 358 t  
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1321 tagagaagag tgtctttgat caactacatt ttatggtttt tatattaa
    
```

//

Revised: May 2, 2000.

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**Nucleotide**

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**Ambrosia artemisiifolia allergen Amb a VI (Amb-a-VI) mRNA, complete cds**

**Views:**

**GenBank view**

Accession: U89793

**Total Bases Sequenced: 357 bp**

**Completed: Apr 1, 1997.**

CDS with gene  
and mRNA

Refresh

gene, tRNA,  
promoter...

FASTA view

### ASN.1 view

### Other features

## Coding Regions

**Legend:**

## Feature table

**Organism:** Ambrosia artemisiifolia

Genetic Code: 1

Lineage: Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; euphyllophytes; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Asteridae; euasterids II; Asterales; Asteraceae; Asteroideae; Heliantheae; Ambrosia.

## Cloning and Expression of Ragweed Allergen Amb a VI

Hiller, K.M., Lubahn, B.C. and Klapper, D.G.

Unpublished

## Direct Submission

Hiller, K.M. and Lubahn, B.C.

Submitted (17-FEB-1997) Micro & Immuno, UNC-CH, CB#7290 631 FLOB, Chapel Hill, NC 27599, USA

*Comments and suggestions to: [info@ncbi.nlm.nih.gov]*



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**1 : U89793 . Ambrosia artemisii...[gi:1916291]**

## Protein

```

LOCUS       AAU89793                357 bp            mRNA            PLN            01-APR-1997
DEFINITION  Ambrosia artemisiifolia allergen Amb a VI (Amb-a-VI) mRNA, complete cds.
ACCESSION   U89793
VERSION     U89793.1   GI:1916291
KEYWORDS    .
SOURCE      common ragweed.
  ORGANISM  Ambrosia artemisiifolia
            Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
            euphyllophytes; Spermatophyta; Magnoliophyta; eudicotyledons; core
            eudicots; Asteridae; euasterids II; Asterales; Asteraceae;
            Asteroideae; Heliantheae; Ambrosia.
REFERENCE   1 (bases 1 to 357)
  AUTHORS   Hiller,K.M., Lubahn,B.C. and Klapper,D.G.
  TITLE     Cloning and Expression of Ragweed Allergen Amb a VI
  JOURNAL   Unpublished
REFERENCE   2 (bases 1 to 357)
  AUTHORS   Hiller,K.M. and Lubahn,B.C.
  TITLE     Direct Submission
  JOURNAL   Submitted (17-FEB-1997) Micro & Immuno, UNC-CH, CB#7290 631 FLOB,
            Chapel Hill, NC 27599, USA
FEATURES             Location/Qualifiers
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     CDS                1..357
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                       DFPVADKLNLDCKSLPV"
BASE COUNT      100 a        81 c        81 g        95 t
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    121 gcaggtttct taacgggtca ggagccaagc aaggcgtgct gcacaggagt caacaatctt
    181 aataactcga gaaaaaccaa agctgatcgt gtggccgtct gcaactgtat caaagaattg
    241 acaaaatcga ttgcttacga tccaaaacgt atgcctcttc tgtctacgaa atgtggcggt
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**major fecal allergen Der p I - house-dust mite (*Dermatophagoides pteronyssinus*) (fragments)**

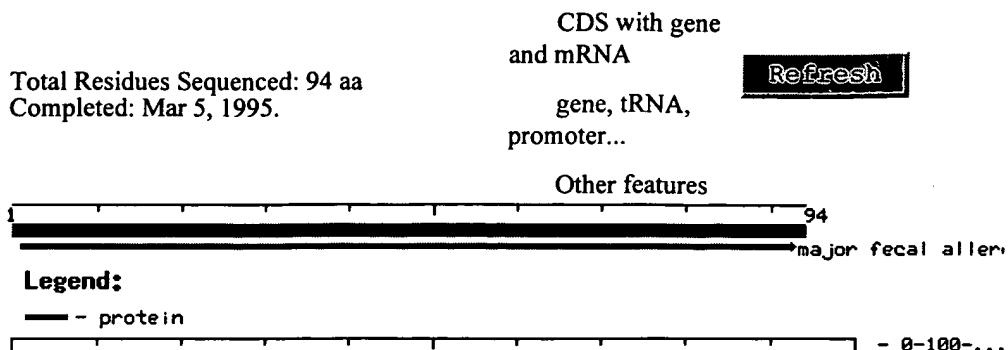
**Views:**

[GenBank view](#)

[FASTA view](#)

[ASN.1 view](#)

[Feature table](#)



Organism: *Dermatophagoides pteronyssinus*

Genetic Code: 1

Lineage: Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Acari; Acariformes; Sarcoptiformes; Astigmata; Analgoidea; Pyroglyphidae; Dermatophagoides.

**Structural studies on the allergen Der p1 from the house dust mite *Dermatophagoides pteronyssinus*: similarity with cysteine proteinases**

Simpson, R.J., Nice, E.C., Moritz, R.L. and Stewart, G.A.

Protein Seq. Data Anal. 2 (1), 17-21 (1989)

89098855

Comments and suggestions to: [\[info@ncbi.nlm.nih.gov\]](mailto:info@ncbi.nlm.nih.gov)





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1 : S03380 . major fecal allerg...[gi:1078971]

PubMed, Related Sequences

LOCUS S03380 94 aa INV 01-SEP-1995  
DEFINITION major fecal allergen Der p I - house-dust mite (Dermatophagoides pteronyssinus) (fragments).  
ACCESSION S03380  
PID g1078971  
VERSION S03380 GI:1078971  
DBSOURCE pir: locus S03380;  
summary: #length 94 #checksum 3067;  
PIR dates: 05-Mar-1995 #sequence\_revision 01-Sep-1995 #text\_change 01-Sep-1995;  
punctuation in sequence.  
KEYWORDS .  
SOURCE Dermatophagoides pteronyssinus.  
ORGANISM Dermatophagoides pteronyssinus  
Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Acari;  
Acariformes; Sarcoptiformes; Astigmata; Analgoidea; Pyroglyphidae;  
Dermatophagoides.  
REFERENCE 1 (residues 1 to 94)  
AUTHORS Simpson,R.J., Nice,E.C., Moritz,R.L. and Stewart,G.A.  
TITLE Structural studies on the allergen Der p1 from the house dust mite Dermatophagoides pteronyssinus: similarity with cysteine proteinases  
JOURNAL Protein Seq. Data Anal. 2 (1), 17-21 (1989)  
MEDLINE 89098855  
FEATURES  
Location/Qualifiers  
source 1..94  
/organism="Dermatophagoides pteronyssinus"  
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Protein 1..94  
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61 fgisnycqiy ppnankdngy qpnxavniv gyxn  
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**Dermatophagoides farinae mRNA for mite allergen Der f II precursor, partial cds, clone:pFL11**

**Views:**

**GenBank view**

Accession: D10449  
Total Bases Sequenced: 485 bp  
Completed: Apr 28, 1993.

CDS with gene  
and mRNA

Refresh

FASTA view

gene, tRNA,  
promoter...

### ASN.1 view

### Other features

## Coding Regions

mitte allergen Der f II pre  
polyA signal

## Feature table

**Legend:**

- - CDS      - - other feature  
 - - 100-200-...

**Organism:** *Dermatophagoides farinae*

Genetic Code: 1

Lineage: Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Acari; Acariformes; Sarcoptiformes; Astigmata; Analgoidea; Pyroglyphidae; Dermatophagoides.

## Cloning and expression of cDNA coding for the major house dust mite allergen Der f II in *Escherichia coli*

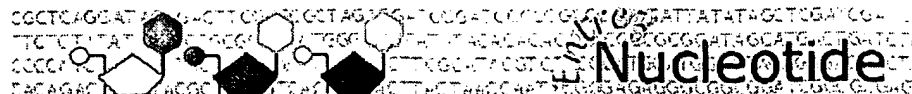
Yuuki,T., Okumura,Y., Ando,T., Yamakawa,H., Suko,M., Haida,M. and Okudaira,H.  
Agric. Biol. Chem. 55 (5), 1233-1238 (1991)  
91291341

## Direct Submission

Yuuki, T.

Submitted (31-JAN-1992) to the DDBJ/EMBL/GenBank databases. Toshifumi Yuuki, Asahi Breweries, Ltd., Central Research Laboratories; 2-13-1, Ohmori-kita, Ohta-ku, Tokyo 143, Japan (Tel:03-5493-3255, Fax:03-5493-7027)

*Comments and suggestions to: [\[info@ncbi.nlm.nih.gov\]](mailto:info@ncbi.nlm.nih.gov)*



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1 : D10449 . Dermatophagoides f...[gi:217307]

PubMed, Protein, Related Sequences

LOCUS DEPDER3 485 bp mRNA INV 01-FEB-2000  
 DEFINITION Dermatophagoides farinae mRNA for mite allergen Der f II precursor partial cds, clone:pFL11.  
 ACCESSION D10449  
 VERSION D10449.1 GI:217307  
 KEYWORDS Der II major allergen group; Der f II; mite allergen.  
 SOURCE Dermatophagoides farinae cDNA to mRNA, clone:pFL11.  
 ORGANISM Dermatophagoides farinae  
 Eukaryota; Metazoa; Arthropoda; Chelicerata; Arachnida; Acari; Acariformes; Sarcoptiformes; Astigmata; Analgoidea; Pyroglyphidae; Dermatophagoides.  
 REFERENCE 1 (bases 1 to 485)  
 AUTHORS Yuuki,T., Okumura,Y., Ando,T., Yamakawa,H., Suko,M., Haida,M. and Okudaira,H.  
 TITLE Cloning and expression of cDNA coding for the major house dust mit allergen Der f II in Escherichia coli  
 JOURNAL Agric. Biol. Chem. 55 (5), 1233-1238 (1991)  
 MEDLINE 91291341  
 REFERENCE 2 (bases 1 to 485)  
 AUTHORS Yuuki,T.  
 TITLE Direct Submission  
 JOURNAL Submitted (31-JAN-1992) to the DDBJ/EMBL/GenBank databases. Toshifumi Yuuki, Asahi Breweries, Ltd., Central Research Laboratories; 2-13-1, Ohmori-kita, Ohta-ku, Tokyo 143, Japan (Tel:03-5493-3255, Fax:03-5493-7027)  
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 source Location/Qualifiers  
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421 aaaaaaaaaa aaatatgaaa attttcacca acatcgaaca aaattcaata accaaaaatt
481 gaac
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CGCTCAGGATACCTGACTTCCTCCGCTAGTCCATCGGATCCCGGCAAGGATTATAGCTCGATCGATCT  
TTCTCTTATACCGCAATATGGGATATATACACACACATCCCGCGATAGCATGACTGATCTA  
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### Human gene for muscarinic acetylcholine receptor HM4

#### Views:

[GenBank view](#)

Accession: [X15266](#)  
Total Bases Sequenced: 1913 bp  
Completed: Feb 19, 1990.

CDS with gene  
and mRNA

[Refresh](#)

gene, tRNA,  
promoter...

[FASTA view](#)

[ASN.1 view](#)

Other features

[Coding Regions](#)

#### Legend:

— CDS

[Feature table](#)

— 1000-2000-...

Organism: [Homo sapiens](#)

Genetic Code: [1](#)

Lineage: Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria;  
Primates; Catarrhini; Hominidae; Homo.

#### Distinct primary structures, ligand-binding properties and tissue-specific expression of four human muscarinic acetylcholine receptors

Peralta, E.G., Ashkenazi, A., Winslow, J.W., Smith, D.H., Ramachandran, J. and Capon, D.J.  
EMBO J. 6 (13), 3923-3929 (1987)

88166632

COMMENT See X15263-15266 for other human muscarinic acetylcholine receptor genes.

Comments and suggestions to: [\[info@ncbi.nlm.nih.gov\]](mailto:info@ncbi.nlm.nih.gov)



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**1 : X15266 . Human gene for mus...**[\[gi:32323\]](#)

PubMed, Protein, Related Sequences, LinkOut

LOCUS	HSHM4	1913 bp	DNA	PRI	14-JAN-2000
DEFINITION	Human gene for muscarinic acetylcholine receptor HM4.				
ACCESSION	X15266 X13530				
VERSION	X15266.1 GI:32323				
KEYWORDS	muscarinic acetylcholine receptor.				
SOURCE	human.				
ORGANISM	<u>Homo sapiens</u> Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.				
REFERENCE	1 (bases 1 to 1913)				
AUTHORS	Peralta,E.G., Ashkenazi,A., Winslow,J.W., Smith,D.H., Ramachandran,J. and Capon,D.J.				
TITLE	Distinct primary structures, ligand-binding properties and tissue-specific expression of four human muscarinic acetylcholine receptors				
JOURNAL	EMBO J. 6 (13), 3923-3929 (1987)				
MEDLINE	<u>88166632</u>				
COMMENT	See X15263-15266 for other human muscarinic acetylcholine receptor genes.				

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BASE COUNT	496 a	529 c	444 g	444 t
ORIGIN				

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Revised: May 2, 2000.

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BLAST PubMed Nucleotide Protein Genome Structure PopSet Taxonomy Help  
**collagen alpha 3(IV) chain precursor, long splice form - human**

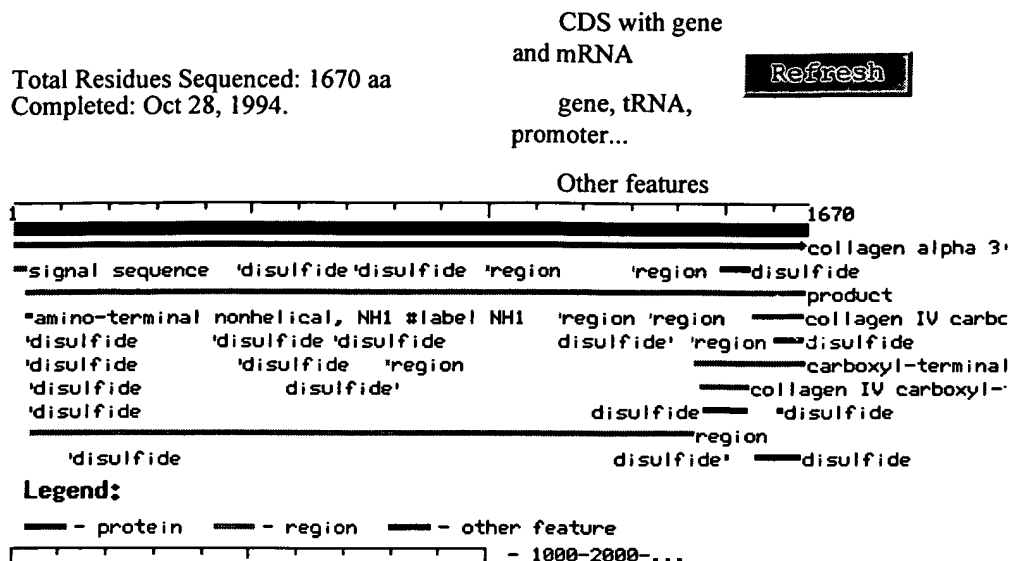
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GenBank view

FASTA view

ASN.1 view

Feature table

Organism: Homo sapiensGenetic Code: 1

Lineage: Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

**Sequence and localization of a partial cDNA encoding the human alpha 3 chain of type IV collagen**

Morrison, K.E., Mariyama, M., Yang-Feng, T.L. and Reeder, S.T.  
 Am. J. Hum. Genet. 49 (3), 545-554 (1991)  
[91353570](#)

**Molecular cloning of the human Goodpasture antigen demonstrates it to be the alpha 3 chain of type IV collagen**

Turner, N., Mason, P.J., Brown, R., Fox, M., Povey, S., Rees, A. and Pusey, C.D.  
 J. Clin. Invest. 89 (2), 592-601 (1992)  
[92147878](#)

**Exon/intron structure of the human alpha 3(IV) gene encompassing the Goodpasture antigen (alpha 3(IV)NC1). Identification of a potentially antigenic region at the triple helix/NC1 domain junction**

Quinones, S., Bernal, D., Garcia-Sogo, M., Elena, S.F. and Saus, J.  
 J. Biol. Chem. 267 (28), 19780-19784 (1992)  
[93015826](#)

**The human mRNA encoding the Goodpasture antigen is alternatively spliced**

Bernal, D., Quinones, S. and Saus, J.  
 J. Biol. Chem. 268 (16), 12090-12094 (1993)  
[93280184](#)

**Exon/intron structure of the human alpha 3(IV) gene encompassing the Goodpasture antigen (alpha 3(IV)NC1). Identification of a potentially antigenic region at the triple helix/NC1 domain junction**

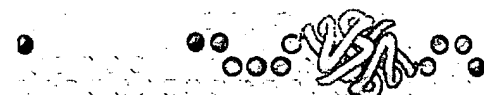
Quinones,S., Bernal,D., Garcia-Sogo,M., Elena,S.F. and Saus,J.  
J. Biol. Chem. 269 (25), 17358 (1994)  
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**Complete primary structure of the human alpha 3(IV) collagen chain. Coexpression of the alpha 3(IV) and alpha 4(IV) collagen chains in human tissues**  
Mariyama,M., Leinonen,A., Mochizuki,T., Tryggvason,K. and Reeders,S.T.  
J. Biol. Chem. 269 (37), 23013-23017 (1994)  
94364994

**COMMENT** Prolines and lysines at the third position of the tripeptide repeating unit (G-X-Y) are hydroxylated to varying extents. Prolines are predominately 4-hydroxylated. Lysines are 5-hydroxylated and subsequently O-glycosylated. In Goodpasture's syndrome, an autoimmune response develops against an epitope in the carboxyl-terminal nonhelical NC1 domain. This minor type IV collagen is thought to form a heterotrimer of two alpha 3(IV) chains and one alpha 4(IV) chain (see PIR:CGHU1B). A polymeric network forms with tetrameric associations among trimer amino-terminal domains (with disulfide and desmosine cross-links), dimeric associations among trimer carboxyl-terminal domains (with disulfide bonds), and both intra-trimer and inter-trimer associations in the interrupted helical domain (with disulfide and desmosine cross-links).

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*Comments and suggestions to: [\[info@ncbi.nlm.nih.gov\]](mailto:info@ncbi.nlm.nih.gov)*



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[Structure](#)
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[History](#)
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# 1 : CGHU3B . collagen alpha 3(I...[gi:1360672]

PubMed, Related Sequences

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 1488/1; 1547/2; 1585/3; 1643/2 #note the alpha 3(IV) and alpha  
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 promotor regions; defects in this gene can result in recessive for  
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 PIR dates: 28-Oct-1994 #sequence\_revision 03-Oct-1995 #text\_change  
 22-Jun-1999.

KEYWORDS alternative splicing; basement membrane; cell binding; coiled coil  
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 trimer; triple helix.

SOURCE human.  
 ORGANISM Homo sapiens  
 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia;  
 Eutheria; Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (residues 1 to 1670)  
 AUTHORS Morrison,K.E., Mariyama,M., Yang-Feng,T.L. and Reeders,S.T.  
 TITLE Sequence and localization of a partial cDNA encoding the human  
 alpha 3 chain of type IV collagen  
 JOURNAL Am. J. Hum. Genet. 49 (3), 545-554 (1991)  
 MEDLINE 91353570

REFERENCE 2 (residues 1 to 1670)  
 AUTHORS Turner,N., Mason,P.J., Brown,R., Fox,M., Povey,S., Rees,A. and  
 Pusey,C.D.  
 TITLE Molecular cloning of the human Goodpasture antigen demonstrates it  
 to be the alpha 3 chain of type IV collagen  
 JOURNAL J. Clin. Invest. 89 (2), 592-601 (1992)  
 MEDLINE 92147878

REFERENCE 3 (residues 1 to 1670)  
 AUTHORS Quinones,S., Bernal,D., Garcia-Sogo,M., Elena,S.F. and Saus,J.  
 TITLE Exon/intron structure of the human alpha 3(IV) gene encompassing  
 the Goodpasture antigen (alpha 3(IV)NC1). Identification of a  
 potentially antigenic region at the triple helix/NC1 domain  
 junction  
 JOURNAL J. Biol. Chem. 267 (28), 19780-19784 (1992)  
 MEDLINE 93015826

REMARK Erratum:[published erratum appears in J Biol Chem 1994 Jun  
 24;269(25):17358]]

REFERENCE 4 (residues 1 to 1670)  
 AUTHORS Bernal,D., Quinones,S. and Saus,J.

TITLE The human mRNA encoding the Goodpasture antigen is alternatively spliced  
 JOURNAL J. Biol. Chem. 268 (16), 12090-12094 (1993)  
 MEDLINE 93280184  
 REFERENCE 5 (residues 1 to 1670)  
 AUTHORS Quinones,S., Bernal,D., Garcia-Sogo,M., Elena,S.F. and Saus,J.  
 TITLE Exon/intron structure of the human alpha 3(IV) gene encompassing the Goodpasture antigen (alpha 3(IV)NC1). Identification of a potentially antigenic region at the triple helix/NC1 domain junction  
 JOURNAL J. Biol. Chem. 269 (25), 17358 (1994)  
 MEDLINE 94274734  
 REMARK annotation; erratum; correction to intronic sequence in A44043  
 REFERENCE 6 (residues 1 to 1670)  
 AUTHORS Mariyama,M., Leinonen,A., Mochizuki,T., Tryggvason,K. and Reeders,S.T.  
 TITLE Complete primary structure of the human alpha 3(IV) collagen chain Coexpression of the alpha 3(IV) and alpha 4(IV) collagen chains in human tissues  
 JOURNAL J. Biol. Chem. 269 (37), 23013-23017 (1994)  
 MEDLINE 94364994  
 COMMENT Prolines and lysines at the third position of the tripeptide repeating unit (G-X-Y) are hydroxylated to varying extents. Prolines are predominately 4-hydroxylated. Lysines are 5-hydroxylated and subsequently O-glycosylated. In Goodpasture's syndrome, an autoimmune response develops against an epitope in the carboxyl-terminal nonhelical NC1 domain. This minor type IV collagen is thought to form a heterotrimer of two alpha 3(IV) chains and one alpha 4(IV) chain (see PIR:CGHU1B). A polymeric network forms with tetrameric associations among trimer amino-terminal domains (with disulfide and desmosine cross-links), dimeric associations among trimer carboxyl-terminal domains (with disulfide bonds), and both intra-trimer and inter-trimer associations in the interrupted helical domain (with disulfide and desmosine cross-links).  
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BLAST PubMed Nucleotide Protein Genome Structure PopSet Taxonomy Help

**Ku autoantigen p70 subunit [human, mRNA, 2123 nt]**

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GenBank view

Accession: [S38729](#)  
Total Bases Sequenced: 2123 bp  
Completed: May 8, 1993.

CDS with gene  
and mRNA

[Refresh](#)

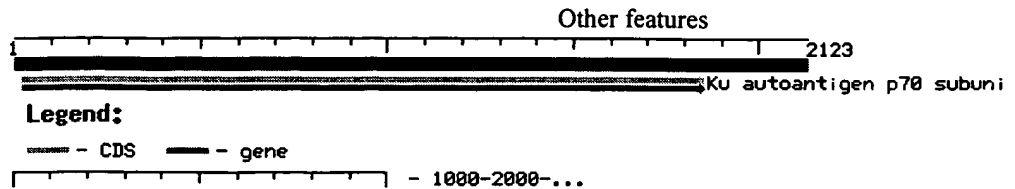
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FASTA view

ASN.1 view

Coding Regions

Feature table



Links:

LocusLink



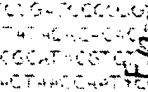



Organism: [Homo sapiens](#)

Genetic Code: [1](#)

Lineage: Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria;  
Primates; Catarrhini; Hominidae; Homo.

**Nucleotide sequence and genomic structure analyses of the p70 subunit of the human Ku autoantigen: evidence for a family of genes encoding Ku (p70)-related polypeptides**  
Griffith,A.J., Craft,J., Evans,J., Mimori,T. and Hardin,J.A.  
Mol. Biol. Rep. 16 (2), 91-97 (1992)  
[92301477](#)

Comments and suggestions to: [\[info@ncbi.nlm.nih.gov\]](mailto:info@ncbi.nlm.nih.gov)

PubMed    Nucleotide    Protein    Genome    Structure    PopSet

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1 : S38729 . Ku autoantigen p70...[gi:250496]

PubMed, Protein, Related Sequences, LinkOut

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 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia;  
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REFERENCE 1 (bases 1 to 2123)  
 AUTHORS Griffith,A.J., Craft,J., Evans,J., Mimori,T. and Hardin,J.A.  
 TITLE Nucleotide sequence and genomic structure analyses of the p70  
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 encoding Ku (p70)-related polypeptides  
 JOURNAL Mol. Biol. Rep. 16 (2), 91-97 (1992)  
 MEDLINE 92301477  
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Revised: May 2, 2000.

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BLAST PubMed Nucleotide Protein Genome Structure PopSet Taxonomy Help

**Homo sapiens myelin-associated oligodendrocyte basic protein (MOBP) mRNA**

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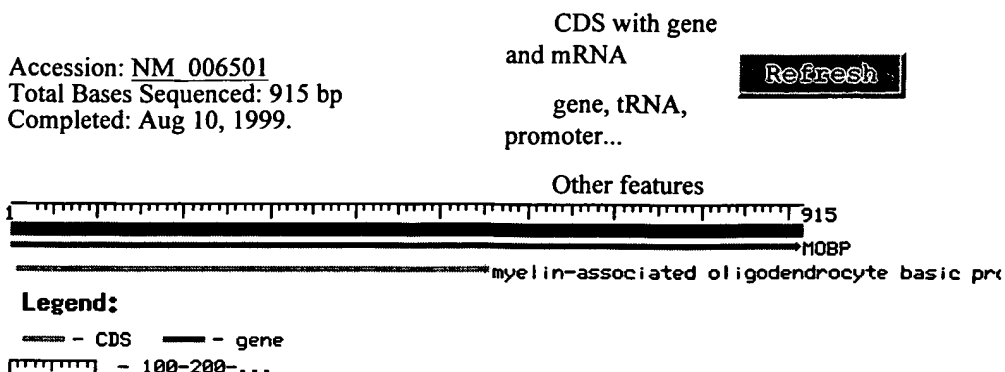
GenBank view

FASTA view

ASN.1 view

Coding Regions

Feature table



Organism: Homo sapiens

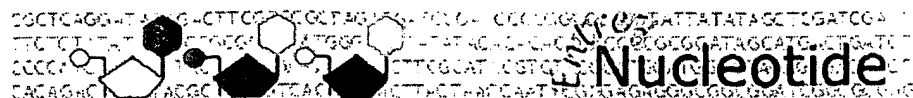
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Lineage: Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

**Cloning and expression of myelin-associated oligodendrocytic basic protein. A novel basic protein constituting the central nervous system myelin**  
Yamamoto, Y., Mizuno, R., Nishimura, T., Ogawa, Y., Yoshikawa, H., Fujimura, H., Adachi, E., Kishimoto, T., Yanagihara, T. and Sakoda, S.  
J. Biol. Chem. 269 (50), 31725-31730 (1994)  
95081123

**COMMENT** REFSEQ: This reference sequence was derived from D28113.1.  
PROVISIONAL RefSeq: This is a provisional reference sequence record that has not yet been subject to human review. The final curated reference sequence record may be somewhat different from this one.

Comments and suggestions to: [\[info@ncbi.nlm.nih.gov\]](mailto:info@ncbi.nlm.nih.gov)



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1 : [NM\\_006501](#) . Homo sapiens myeli...[gi:5729930][PubMed](#), [Protein](#), [Related Sequences](#), [LinkOut](#)

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 VERSION NM\_006501.1 GI:5729930  
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 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia;  
 Eutheria; Primates; Catarrhini; Hominidae; Homo.  
 REFERENCE 1 (sites)  
 AUTHORS Yamamoto,Y., Mizuno,R., Nishimura,T., Ogawa,Y., Yoshikawa,H.,  
 Fujimura,H., Adachi,E., Kishimoto,T., Yanagihara,T. and Sakoda,S.  
 TITLE Cloning and expression of myelin-associated oligodendrocytic basic  
 protein. A novel basic protein constituting the central nervous  
 system myelin  
 JOURNAL J. Biol. Chem. 269 (50), 31725-31730 (1994)  
 MEDLINE 95081123  
 COMMENT REFSEQ: This reference sequence was derived from D28113.1.  
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**Nucleotide**

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**Views:**

**GenBank view**

Accession: X05615  
Total Bases Sequenced: 8448 bp  
Completed: Apr 2, 1988.

CDS with gene  
and mRNA

gene, tRNA,  
promoter...

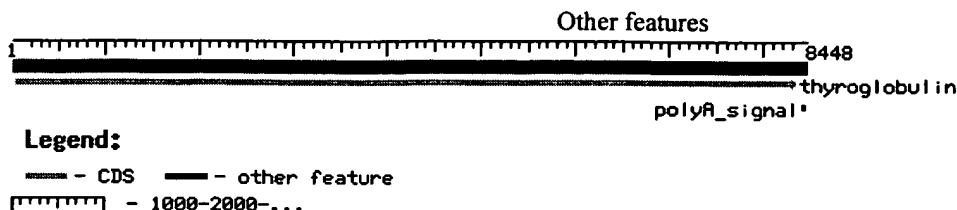
Refresh

FASTA view

### ASN.1 view

## Coding Regions

## Feature table



**Organism:** *Homo sapiens*

Genetic Code: 1

Lineage: Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

### Primary structure of human thyroglobulin deduced from the sequence of its 8448-base complementary DNA

Malthiery, Y. and Lissitzky, S.

Eur. J. Biochem. 165 (3), 491-498 (1987)

87246630

### Direct Submission

Malthiery, Y.

Submitted (07-APR-1988) to the EMBL/GenBank/DDBJ databases

### A major human thyroglobulin epitope defined with monoclonal antibodies is mainly recognized by human autoantibodies


Henry, M., Zanelli, E., Piechaczyk, M., Pau, B. and Malthiery, Y.

Eur. J. Immunol. 22 (2), 315-319 (1992)

92164705

**COMMENT** patient); Data kindly reviewed (07-APR-1988) by Malthiery Y.

*Comments and suggestions to: [info@ncbi.nlm.nih.gov]*



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PubMed Nucleotide Protein Genome Structure PopSet

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1 : X05615 . Human mRNA for thy...[gi:37173]

PubMed, Protein, Related Sequences

LOCUS HSTHYRR 8448 bp mRNA PRI 17-FEB-1997

DEFINITION Human mRNA for thyroglobulin.

ACCESSION X05615

VERSION X05615.1 GI:37173

KEYWORDS thyroglobulin.

SOURCE human.

ORGANISM Homo sapiens  
 Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria;  
 Primates; Catarrhini; Hominidae; Homo.

REFERENCE 1 (bases 1 to 8446)

AUTHORS Malthiery, Y. and Lissitzky, S.

TITLE Primary structure of human thyroglobulin deduced from the sequence of its 8448-base complementary DNA

JOURNAL Eur. J. Biochem. 165 (3), 491-498 (1987)

MEDLINE 87246630

REFERENCE 2 (bases 1 to 8448)

AUTHORS Malthiery, Y.

TITLE Direct Submission

JOURNAL Submitted (07-APR-1988) to the EMBL/GenBank/DDBJ databases

REFERENCE 3 (bases 1 to 8448)

AUTHORS Henry, M., Zanelli, E., Piechaczyk, M., Pau, B. and Malthiery, Y.

TITLE A major human thyroglobulin epitope defined with monoclonal antibodies is mainly recognized by human autoantibodies

JOURNAL Eur. J. Immunol. 22 (2), 315-319 (1992)

MEDLINE 92164705

COMMENT patient);  
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**H.sapiens mRNA for put. B7,3 molecule of CD80-CD60 protein family**

### Views:

**GenBank view**

FASTA view

### ASN.1 view

## Coding Regions

## Feature table

Accession: Y07827  
Total Bases Sequenced: 1182 bp  
Completed: Jan 8, 1997.

CDS with gene  
and mRNA

gene, tRNA,  
promoter...

Refresh

## Other features

1182  
put. B7,3 molecule

**Legend:**

~~100-200-...~~ - CDS  
100-200-...

Organism: *Homo sapiens*

Genetic Code: 1

Lineage: Eukaryota; Metazoa; Chordata; Vertebrata; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

## **Cloning, structural analysis and mapping of B30 and B7 family members, to the MHC and other chromosomal regions. Toward the identification of the ancestral major histocompatibility complex**

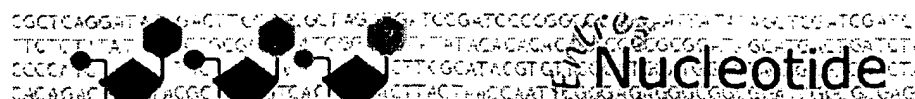
Henry,J., Ribouchon,M.T., Depetris,D., Mattei,M.G., Offer,C., Tazi-Ahnini,R. and Pantarotti,P.  
Unpublished

## Direct Submission

Pontarotti, P.

Submitted (06-SEP-1996) P. Pontarotti, Unite 119 INSERM, 27 bd.Lei Roure, 13009  
Marseille, FRANCE

*Comments and suggestions to: [\[info@ncbi.nlm.nih.gov\]](mailto:info@ncbi.nlm.nih.gov)*



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**1 : Y07827 . H.sapiens mRNA for...[gi:1770367]**

Protein, Related Sequences, LinkOut

```

LOCUS       HSB73                1182 bp      mRNA                PRI      08-JAN-1997
DEFINITION  H.sapiens mRNA for put. B7,3 molecule of CD80-CD60 protein family.
ACCESSION   Y07827
VERSION     Y07827.1   GI:1770367
KEYWORDS    major histocompatibility complex.
SOURCE      human.
  ORGANISM  Homo sapiens
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            Primates; Catarrhini; Hominidae; Homo.
REFERENCE   1 (bases 1 to 1182)
  AUTHORS   Henry,J., Ribouchon,M.T., Depetris,D., Mattei,M.G., Offer,C.,
            Tazi-Ahnini,R. and Pantarotti,P.
  TITLE     Cloning, structural analysis and mapping of B30 and B7 family
            members, to the MHC and other chromosomal regions. Toward the
            identification of the ancestral major histocompatibility complex
  JOURNAL   Unpublished
REFERENCE   2 (bases 1 to 1182)
  AUTHORS   Pontarotti,P.
  TITLE     Direct Submission
  JOURNAL   Submitted (06-SEP-1996) P. Pontarotti, Unite 119 INSERM, 27 bd.Lei
            Roure, 13009 Marseille, FRANCE
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### B.verrucosa mRNA for pollen allergen Bet v 4

**Views:**

**GenBank view**

Accession: Y12560  
Total Bases Sequenced: 496 bp  
Completed: Apr 24, 1997.

CDS with gene  
and mRNA

Refresh

gene, tRNA,  
promoter...

FASTA view

### Other features

### ASN.1 view



## Coding Regions

**Legend:**

— CDS — gene  
— 100-200-...

## Feature table

**Organism:** *Betula pendula*

**Genetic Code: 1**

Lineage: Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; euphyllophytes; Spermatophyta; Magnoliophyta; eudicotyledons; Rosidae; Fagales; Betulaceae; Betula.

## Molecular characterization, expression in *Escherichia coli*, and epitope analysis of a two EF-hand calcium-binding birch pollen allergen, Bet v 4

Twardosz,A., Hayek,B., Seiberler,S., Vangelista,L., Elfman,L., Gronlund,H., Kraft,D. and Valenta,R.

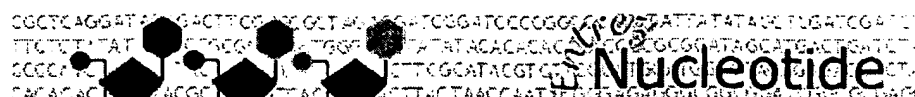
Biochem. Biophys. Res. Commun. 239 (1), 197-204 (1997)  
98005106

## Direct Submission

Valenta, R.

Submitted (14-APR-1997) R. Valenta, Institute of General & Experimental Pathology,  
General Hospital, Waehringer Guertel 18-20, 1090 Vienna, AUSTRIA

*Comments and suggestions to: [info@ncbi.nlm.nih.gov]*



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